PII: S0079-6107(99)00013-9

AUTHOR INDEX TO VOLUME 71

Adams P. D., 135
Arrondo J. L. R., 367

Brunger A. T., 135

Cowtan K. D., 245
Craig K. L., 299

D'Alessio G., 271
Davis B. K., 157

Frishman D., 1

Goñi F. M., 367

Hounsgaard J., 329

Koller F., 19

Mewes H.-W., 1

Rice L. M., 135
Russo R. E., 329

Shimoni Y., 67

Tachibana M., 109
Tyers M., 299

Zámocký M., 19
Zhang K. Y. J., 245

PII: S0079-6107(99)00015-2

AUTHOR INDEX TO VOLUME 72

Ahn N. G., 501
Akamine P., 313

Brette F., 29
Bubeck-Wardenburg J., 373

Calaghan S. C., 59
Cazorla O., 29
Chan A. C., 373
Cobb M. H., 479
Császár A. G., 243

Damen J. E., 423
Duronio V., 423

Fu C., 373
Fütterer K., 373

Hauck C. R., 435
Helgason C. D., 423
Hubbard S. R., 343
Huber M., 423
Humphries R. K., 423
Hunter P., 91

Jaenicke R., 155

Kohl P., 91
Krystal G., 423

Lab M., 3
Lab M. J., 7
Le Guennec J.-Y., 29
Liu L., 423

Madhusudan, 313

Noble D., 1, 91

Pappu R., 373
Pascarel C., 29
Pawson T., 311
Perczel A., 243

Radzio-Andzelm E., 313
Resing K. A., 501

Samelson L. E., 393
Scheid M., 423
Schlaepfer D. D., 435
Shuai K., 405
Sieg D. J., 435
Smith C. M., 313, 525
Songyang Z., 359
Sutton P. M. I., 139

Taggart P., 139
Taylor S. S., 313

Waksman G., 373
Ware M. D., 423
White E., 59
Wong J., 373

PII: S0079-6107(99)00012-7

SUBJECT INDEX TO VOLUME 71

Arrhythmia: 139

C-Src: 435
Cell migration: 435

Electrophysiology: 139
ERK2: 435

FAK: 435

Growth factor receptor: 343

Human heart: 139

Ischaemia: 139

Mechano-electric feedback: 139

Protein tyrosine kinase: 343
Pyk2: 435

Signal transduction: 343

Tyrosine phosphorylation: 343

X-ray crystallography: 343

PII: S0079-6107(99)00016-4

SUBJECT INDEX TO VOLUME 72

Band analysis: 367
Biosynthetic pathways: 157

Catalase-peroxidase: 19
Codon evolution tree: 157
Comparative length: 157

Difference spectroscopy: 367

'Fossil' proteins: 157

Infrared spectroscopy: 367
Intersubunit interfaces: 271
Intersubunit swap: 271

Manganese catalase: 19
Membrane proteins: 367
Molecular evolution: 271

Oligomers: 271
Origin of life: 157
Oxidative stress: 19

Protein folding: 19

Structure-function relationship: 367

tRNA phylogeny: 157
Typical catalase: 19